

## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.**

Application Serial Number: 09/610,551E  
Source: IFU/6  
Date Processed by STIC: 1/7/05

***ENTERED***



IFW/6

## RAW SEQUENCE LISTING

DATE: 11/07/2005

PATENT APPLICATION: US/09/610,551E

TIME: 15:09:46

Input Set : A:\409.1D2.TXT

Output Set: N:\CRF4\11072005\I610551E.raw

4 <110> APPLICANT: Barbas, Carlos F.  
 5 Burton, Dennis R.  
 6 Lerner, Richard A.  
 8 <120> TITLE OF INVENTION: Methods for producing antibody libraries  
 9 using universal or randomized immunoglobulin light chains  
 12 <130> FILE REFERENCE: TSRI 409.1D2  
 14 <140> CURRENT APPLICATION NUMBER: US 09/610,551E  
 15 <141> CURRENT FILING DATE: 2000-07-05  
 17 <150> PRIOR APPLICATION NUMBER: US 08/931,645  
 18 <151> PRIOR FILING DATE: 1997-09-16  
 20 <150> PRIOR APPLICATION NUMBER: US 08/300,386  
 21 <151> PRIOR FILING DATE: 1994-09-02  
 23 <150> PRIOR APPLICATION NUMBER: US 08/174,674  
 24 <151> PRIOR FILING DATE: 1993-12-28  
 26 <150> PRIOR APPLICATION NUMBER: US 07/826,623  
 27 <151> PRIOR FILING DATE: 1992-01-27  
 29 <160> NUMBER OF SEQ ID NOS: 74  
 31 <170> SOFTWARE: FastSEQ for Windows Version 4.0  
 33 <210> SEQ ID NO: 1  
 34 <211> LENGTH: 687  
 35 <212> TYPE: DNA  
 36 <213> ORGANISM: Artificial Sequence  
 38 <220> FEATURE:  
 39 <223> OTHER INFORMATION: Synthesized  
 41 <400> SEQUENCE: 1  
 42 ctcgagcagt ctggggctga ggtgaagaag cctgggtcct cggtaaggt ctcctgcagg 60  
 43 gcttctggag gcaccaa caattatgcc atcagctgg tgcgacaggc ccctggacaa 120  
 44 gggcttgagt ggatgggagg gatctccct ttccgtataa cagcaaagta cgacacaacac 180  
 45 ttccaggcga gagtccat taccgcggac gaatccacgg gcacagccta catggagctg 240  
 46 agcagctga gatctgagga cacggccata tattattgtg cgagagggga tacgatttt 300  
 47 ggagtgacca tggatacta cgctatggac gtctggggcc aagggaccac ggtcaccgtc 360  
 48 tccgcagcct ccaccaaggg cccatcggtc ttccccctgg caccctcctc caagagcacc 420  
 49 tctggggca cagcggccct gggctgcctg gtcaaggact acttccccga accggtgacg 480  
 50 gtgtcgatggaa actcaggcgc cctgaccaggc ggctgcaca cttcccgcc tgcctacag 540  
 51 tcctcaggac tctactccct cagcagcgtg gtgaccgtgc cttccagcag cttgggcacc 600  
 52 cagacccata tctgcaacgt gaatcacaag cccagcaaca ccaaggtgga caagaaagca 660  
 53 gagcccaaat ctgtgacaa aactagt 687  
 55 <210> SEQ ID NO: 2  
 56 <211> LENGTH: 646  
 57 <212> TYPE: DNA  
 58 <213> ORGANISM: Artificial Sequence  
 60 <220> FEATURE:  
 61 <223> OTHER INFORMATION: Synthesized

R6

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63 <400> SEQUENCE: 2  
 64 gagctcacgc agtctccagg caccctgtct ttgtctccag gggaaagagc caccctctcc 60  
 65 tgcagggcca gtcacagtgt tagcagggcc tacttagct ggtaccagca gaaacctggc 120  
 66 caggctcca ggctcctcat ctatggtaca tccagcaggg ccactggcat cccagacagg 180  
 67 tccagtgca gtgggtctgg gacagacttc actctcacca tcagcagact ggagcctgaa 240  
 68 gattttcag tgtaactactg tcagcagttat ggtggctcac cgtggttcgg ccaagggacc 300  
 69 aagggtgaac tcaaacgaac tggctgcga ccatctgtct tcatttccc gccatctgat 360  
 70 gaggcgttga aatctggaaac tgcctctgtt gtgtgcctgc tgaataactt ctatcccaga 420  
 71 gaggccaaag tacagtggaa ggtggataac gcccctcaat cgggtaactc ccaggagagt 480  
 72 gtcacagagc aggacagcaa ggacagcacc tacagcctca gcagcaccct gacgctgagc 540  
 73 aaagcagact acgagaaaca caaagtctac gcctgcgaag tcacccatca gggcctgagt 600  
 74 tcgcccgtca caaagagctt caacagggga gagtgttaat tctaga 646  
 76 <210> SEQ ID NO: 3  
 77 <211> LENGTH: 21  
 78 <212> TYPE: DNA  
 79 <213> ORGANISM: Artificial Sequence  
 81 <220> FEATURE:  
 82 <223> OTHER INFORMATION: Synthesized  
 84 <400> SEQUENCE: 3  
 85 gaattctaaa ctagcttagtc g 21  
 87 <210> SEQ ID NO: 4  
 88 <211> LENGTH: 21  
 89 <212> TYPE: DNA  
 90 <213> ORGANISM: Artificial Sequence  
 92 <220> FEATURE:  
 93 <223> OTHER INFORMATION: Synthesized  
 95 <400> SEQUENCE: 4  
 96 atactgctga cagtaataaca c 21  
 98 <210> SEQ ID NO: 5  
 99 <211> LENGTH: 57  
 100 <212> TYPE: DNA  
 101 <213> ORGANISM: Artificial Sequence  
 103 <220> FEATURE:  
 104 <223> OTHER INFORMATION: Synthesized  
 -> 106 <221> NAME/KEY: modified\_base  
 107 <222> LOCATION: 19, 20, 22, 23, 25, 26, 28, 29  
 108 <223> OTHER INFORMATION: n = G, A, T, or C  
 -> 110 <221> modified\_base  
 111 <222> LOCATION: 21, 24, 27, 30  
 112 <223> OTHER INFORMATION: k = G or T  
 -> 114 <400> 5  
 -> 115 tattactgtc agcagtatnn knnknnknnk actttcggcg gagggaccaa ggtggag 57  
 117 <210> SEQ ID NO: 6  
 118 <211> LENGTH: 21  
 119 <212> TYPE: DNA  
 120 <213> ORGANISM: Artificial Sequence  
 122 <220> FEATURE:  
 123 <223> OTHER INFORMATION: Synthesized  
 125 <400> SEQUENCE: 6

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126 aatacgactc actataggc g 21  
 128 <210> SEQ ID NO: 7  
 129 <211> LENGTH: 48  
 130 <212> TYPE: DNA  
 131 <213> ORGANISM: Artificial Sequence  
 133 <220> FEATURE:  
 134 <223> OTHER INFORMATION: Synthesized  
 -> 136 <221> NAME/KEY: modified base  
 137 <222> LOCATION: 19, 20, 22, 23, 25, 26, 28, 29  
 138 <223> OTHER INFORMATION: n = G, A, T, or C  
 -> 140 <221> modified base  
 141 <222> LOCATION: 21, 24, 27, 30  
 142 <223> OTHER INFORMATION: k = G or T  
 -> 144 <400> 7  
 -> 145 tattactgtc agcagtatnn knnknnknnk actttcggcg gagggacc 48  
 147 <210> SEQ ID NO: 8  
 148 <211> LENGTH: 60  
 149 <212> TYPE: DNA  
 150 <213> ORGANISM: Artificial Sequence  
 152 <220> FEATURE:  
 153 <223> OTHER INFORMATION: Synthesized  
 -> 155 <221> NAME/KEY: modified base  
 156 <222> LOCATION: 19, 20, 22, 23, 25, 26, 28, 29, 31, 32  
 157 <223> OTHER INFORMATION: n = G, A, T, or C  
 -> 159 <221> modified base  
 160 <222> LOCATION: 21, 24, 27, 30, 33  
 161 <223> OTHER INFORMATION: k = G or T  
 -> 163 <400> 8  
 -> 164 tattactgtc agcagtatnn knnknnknnk nnkactttcg gcggagggac caaggtggag 60  
 167 <210> SEQ ID NO: 9  
 168 <211> LENGTH: 51  
 169 <212> TYPE: DNA  
 170 <213> ORGANISM: Artificial Sequence  
 172 <220> FEATURE:  
 173 <223> OTHER INFORMATION: Synthesized  
 -> 175 <221> NAME/KEY: modified base  
 176 <222> LOCATION: 19, 20, 22, 23, 25, 26, 28, 29, 31, 32  
 177 <223> OTHER INFORMATION: n = G, A, T, or C  
 -> 179 <221> modified base  
 180 <222> LOCATION: 21, 24, 27, 30, 33  
 181 <223> OTHER INFORMATION: k = G or T  
 -> 183 <400> 9  
 -> 184 tattactgtc agcagtatnn knnknnknnk nnkactttcg gcggagggac c 51  
 186 <210> SEQ ID NO: 10  
 187 <211> LENGTH: 75  
 188 <212> TYPE: DNA  
 189 <213> ORGANISM: Artificial Sequence  
 191 <220> FEATURE:  
 192 <223> OTHER INFORMATION: Synthesized

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-> 194 <221> NAME/KEY: modified_base
195 <222> LOCATION: 31, 32, 34, 35, 37, 38, 40, 41, 43, 44, 46, 47
196 <223> OTHER INFORMATION: n = G, A, T, or C
-> 198 <221> modified_base
199 <222> LOCATION: 33, 36, 39, 42, 45, 48
200 <223> OTHER INFORMATION: k = G or T
-> 202 <400> 10
-> 203 gatttgcag tgtattactg tcagcagtat nnknnknnkn nknnknnkac tttcggcgga 60
204 gggaccaagg tggag 75
206 <210> SEQ ID NO: 11
207 <211> LENGTH: 54
208 <212> TYPE: DNA
209 <213> ORGANISM: Artificial Sequence
211 <220> FEATURE:
212 <223> OTHER INFORMATION: Synthesized
-> 214 <221> NAME/KEY: modified_base
215 <222> LOCATION: 19, 20, 22, 23, 25, 26, 28, 29, 31, 32, 34, 35
216 <223> OTHER INFORMATION: n = G, A, T, or C
-> 218 <221> modified_base
219 <222> LOCATION: 21, 24, 27, 30, 33, 36
220 <223> OTHER INFORMATION: k = G or T
-> 222 <400> 11
-> 223 tattactgtc agcagtatnn knnknnknnk nnknnkactt tcggcggagg gacc 54
225 <210> SEQ ID NO: 12
226 <211> LENGTH: 75
227 <212> TYPE: DNA
228 <213> ORGANISM: Artificial Sequence
230 <220> FEATURE:
231 <223> OTHER INFORMATION: Synthesized
-> 233 <221> NAME/KEY: modified_base
234 <222> LOCATION: 22, 23, 25, 26, 28, 29, 31, 32, 34, 35, 37, 38, 40, 41, 43,
235 44, 46, 47, 49, 50
236 <223> OTHER INFORMATION: n = G, A, T, or C
-> 238 <221> modified_base
239 <222> LOCATION: 24, 27, 30, 33, 36, 39, 42, 45, 48, 51
240 <223> OTHER INFORMATION: k = G or T
-> 242 <400> 12
-> 243 gatttgcag tgtattactg tnnknnknnk nnknnknnkn nknnknnknn kttcggcgga 60
244 gggaccaagg tggag 75
246 <210> SEQ ID NO: 13
247 <211> LENGTH: 70
248 <212> TYPE: DNA
249 <213> ORGANISM: Artificial Sequence
251 <220> FEATURE:
252 <223> OTHER INFORMATION: Synthesized
-> 254 <221> NAME/KEY: modified_base
255 <222> LOCATION: 26, 29, 32, 35, 38, 41, 44, 47
256 <223> OTHER INFORMATION: m = A or C
-> 258 <221> modified_base

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## RAW SEQUENCE LISTING

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259 <222> LOCATION: 27, 28, 30, 31, 33, 34, 36, 37, 39, 40, 42, 43, 45, 46, 48,  
 260 49

261 <223> OTHER INFORMATION: n = G, A, T, or C

-> 263 <400> 13

-> 264 gttccacatt ggtcccttgg ccgaamnnmn nmnnnnnnnn mnmmnnnnna cagtagtaca 60  
 265 ctgcaaaatc 70

267 <210> SEQ ID NO: 14

268 <211> LENGTH: 76

269 <212> TYPE: DNA

270 <213> ORGANISM: Artificial Sequence

272 <220> FEATURE:

273 <223> OTHER INFORMATION: Synthesized

-> 275 <221> NAME/KEY: modified\_base

276 <222> LOCATION: 26, 29, 32, 35, 38, 41, 44, 47, 50, 53

277 <223> OTHER INFORMATION: m = A or C

-> 279 <221> modified\_base

280 <222> LOCATION: 27, 28, 30, 31, 33, 34, 36, 37, 39, 40, 42, 43, 45, 46, 48,  
 281 49, 51, 52, 54, 55

282 <223> OTHER INFORMATION: n = G, A, T, or C

-> 284 <400> 14

-> 285 gttccacatt ggtcccttgg ccgaamnnmn nmnnnnnnnn mnmmnnnnnm nmnnnacagt 60  
 286 agtacactgc aaaatc 76

288 <210> SEQ ID NO: 15

289 <211> LENGTH: 94

290 <212> TYPE: DNA

291 <213> ORGANISM: Artificial Sequence

293 <220> FEATURE:

294 <223> OTHER INFORMATION: Synthesized

-> 296 <221> NAME/KEY: modified\_base

297 <222> LOCATION: 26, 29, 32, 35, 38, 41, 44, 47, 50, 53, 56, 59, 62, 65, 68,  
 298 71

299 <223> OTHER INFORMATION: m = A or C

-> 301 <221> modified\_base

302 <222> LOCATION: 27, 28, 30, 31, 33, 34, 36, 37, 39, 40, 42, 43, 45, 46, 48,  
 303 49, 51, 52, 54, 55, 57, 58, 60, 61, 63, 64, 66, 67, 69,  
 304 70, 72, 73

305 <223> OTHER INFORMATION: n = G, A, T, or C

-> 307 <400> 15

-> 308 gttccacatt ggtcccttgg ccgaamnnmn nmnnnnnnnn mnmmnnnnnm nmnnnnnnmn 60

-> 309 nmnnnnnnnn mnnaacagt tacactgcaa aatc 94

311 <210> SEQ ID NO: 16

312 <211> LENGTH: 25

313 <212> TYPE: DNA

314 <213> ORGANISM: Artificial Sequence

316 <220> FEATURE:

317 <223> OTHER INFORMATION: Synthesized

319 <400> SEQUENCE: 16

320 ttcggccaag ggaccaaggt ggaac 25

322 <210> SEQ ID NO: 17

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 11/07/2005  
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Input Set : A:\409.1D2.TXT  
Output Set: N:\CRF4\11072005\1610551E.raw

**base Note:**

**3 of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> <223> fields of each sequence which presents at least one n or Xaa.**

1#:5; N Pos. 19,20,22,23,25,26,28,29  
1#:7; N Pos. 19,20,22,23,25,26,28,29  
1#:8; N Pos. 19,20,22,23,25,26,28,29,31,32  
1#:9; N Pos. 19,20,22,23,25,26,28,29,31,32  
1#:10; N Pos. 31,32,34,35,37,38,40,41,43,44,46,47  
1#:11; N Pos. 19,20,22,23,25,26,28,29,31,32,34,35  
1#:12; N Pos. 22,23,25,26,28,29,31,32,34,35,37,38,40,41,43,44,46,47,49,50  
1#:13; N Pos. 27,28,30,31,33,34,36,37,39,40,42,43,45,46,48,49  
1#:14; N Pos. 27,28,30,31,33,34,36,37,39,40,42,43,45,46,48,49,51,52,54,55  
1#:15; N Pos. 27,28,30,31,33,34,36,37,39,40,42,43,45,46,48,49,51,52,54,55  
1#:15; N Pos. 57,58,60,61,63,64,66,67,69,70,72,73  
1#:19; N Pos. 22,23,25,26,28,29,34,35  
1#:21; N Pos. 25,26,28,29,31,32,34,35,37,38,40,41,43,44,49,50  
1#:22; N Pos. 25,26,28,29,31,32,34,35,37,38,40,41,43,44,46,47,49,50,52,53  
1#:22; N Pos. 55,56,58,59,61,62,67,68  
1#:23; N Pos. 19,20,22,23,25,26,28,29,31,32  
1#:24; N Pos. 19,20,22,23,25,26,28,29,31,32,34,35,37,38,40,41,43,44,46,47  
1#:25; N Pos. 19,20,22,23,25,26,28,29,31,32,34,35,37,38,40,41,43,44,46,47  
1#:25; N Pos. 49,50,52,53,55,56,58,59,61,62,64,65  
1#:73; N Pos. 1,2,4,5,7,8,10,11,13,14,16,17,19,20,22,23,25,26,28,29,31,32  
1#:73; N Pos. 34,35,37,38,40,41,43,44,46,47,49,50,52,53,55,56,58,59,61,62  
1#:73; N Pos. 64,65,67,68,70,71  
1#:74; N Pos. 2,3,5,6,8,9,11,12,14,15,17,18,20,21,23,24,26,27,29,30,32,33  
1#:74; N Pos. 35,36,38,39,41,42,44,45,47,48,50,51,53,54,56,57,59,60,62,63  
1#:74; N Pos. 65,66,68,69,71,72

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Input Set : A:\409.1D2.TXT  
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106 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
 110 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:5  
 114 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:5  
 115 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0  
 136 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
 140 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:7  
 144 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:7  
 145 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0  
 155 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
 159 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8  
 163 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8  
 164 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0  
 175 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
 179 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:9  
 183 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:9  
 184 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0  
 194 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
 198 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:10  
 202 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:10  
 203 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0  
 214 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
 218 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:11  
 222 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:11  
 223 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0  
 233 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
 238 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:12  
 242 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:12  
 243 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0  
 254 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
 258 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:13  
 263 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:13  
 264 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0  
 275 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
 279 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:14  
 284 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:14  
 285 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0  
 296 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
 301 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:15  
 307 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:15  
 308 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0  
 309 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:60  
 352 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
 356 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:19  
 360 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:19  
 361 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0  
 382 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
 386 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:21  
 390 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:21

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391 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0  
102 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
107 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:22  
111 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:22  
112 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0  
113 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:60  
123 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
127 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:23  
131 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:23  
132 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0  
142 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
147 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24  
151 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24  
152 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0  
163 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
169 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:25  
174 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:25  
175 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0  
176 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:60  
189 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
196 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:73  
201 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:73  
206 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:73  
207 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:73 after pos.:0  
208 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:73 after pos.:60  
218 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
223 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:74  
230 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:74  
235 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:74  
236 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:74 after pos.:0  
237 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:74 after pos.:60